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May 22, 2008

Mr. Steve Trent Fluor Hanford Inc. 1200 Jadwin Avenue Richland, WA 99352

Reference:

P.O. #33677

Eberline Services R8-04-057-7075, SDG H3681

Dear Mr. Trent:

Enclosed is the data report for one solid (soil) sample designated under SAF No. F08-043 received at Eberline Services on April 9, 2008. The sample was analyzed according to the accompanying chain-of-custody document.

Please call if you have any questions concerning this report.

Sincerely,

Melissa C. Mannion

Senior Program Manager

nellero Mannion

MCM/njv

Enclosure: Data Package

1.0 GENERAL

Fluor Hanford Inc. (FH) Sample Delivery Group H3681 was composed of one solid (soil) sample designated under SAF No. F08-043 with a Project Designation of: 216-A-30 Crib Sampling.

The sample was received as stated on the Chain-of-Custody document. Any discrepancies are noted on the Eberline Services Sample Receipt Checklist.

2.0 ANALYSIS NOTES

2.1 Tritium Analysis

No problems were encountered during the course of the analyses.

2.2 Nickel-63 Analysis

No problems were encountered during the course of the analyses.

2.3 Technetium-99 Analysis

No problems were encountered during the course of the analyses.

2.4 Isotopic Thorium Analysis

No problems were encountered during the course of the analyses.

3.0 Case Narrative Certification Statement

"I certify that this data package is in compliance with the SOW, both technically and for completeness, for other than the conditions detailed above. Release of the data obtained in this hard copy data package has been authorized by the Laboratory Manager or a designee, as verified by the following signature."

Melissa C. Mannion

Senior Program Manager

Milina Mannes

Date

EBERLINE SERVICES/RICHMOND SAMPLE DELIVERY GROUP H3681

SDG 7075 Contact Melissa C. Mannion

Client <u>Hanford</u> Contract No. 33677 Case no SDG H3681

SUMMARY DATA SECTION

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Prepared by

Melissa Mannion

Reviewed by

SAMPLE DELIVERY GROUP H3681

SDG 7075 Contact Melissa C. Mannion

REPORT GUIDE

Client Hanford Contract No. 33677 Case no SDG H3681

ABOUT THE DATA SUMMARY SECTION

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

SAMPLE SUMMARIES

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

WORK SUMMARY

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

METHOD BLANKS

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

LAB CONTROL SAMPLES

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

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SAMPLE DELIVERY GROUP H3681

SDG 7075
Contact Melissa C. Mannion

GUIDE, cont.

Client <u>Hanford</u>
Contract <u>No. 33677</u>
Case no <u>SDG H3681</u>

ABOUT THE DATA SUMMARY SECTION

DUPLICATES

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

MATRIX SPIKES

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

DATA SHEETS

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

METHOD SUMMARIES

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

REPORT GUIDES

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

REPORT GUIDES
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SUMMARY DATA SECTION
Page 2

SAMPLE DELIVERY GROUP H3681

LAB SAMPLE SUMMARY

Client Hanford
Contract No. 33677
Case no SDG H3681

LAB SAMPLE ID	CLIENT SAMPLE ID	LOCATION	MATRIX	LEVEL	SAF NO	CHAIN OF CUSTODY	COLLECTED
R804057-01	B1TH05	C5941, I-033	SOLID		F08-043	F08-043-150	03/19/08 08:50
R804057-02	Lab Control Sample		SOLID		F08-043		
R804057-03	Method Blank		SOLID		F08-043		
R804057-04	Duplicate (R804057-01)	C5941, I-033	SOLID		F08-043		03/19/08 08:50

LAB SUMMARY

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SUMMARY DATA SECTION

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SDG 7075

Contact Melissa C. Mannion

SAMPLE DELIVERY GROUP H3681

SDG 7075
Contact Melissa C. Mannion

QC SUMMARY

Client Hanford
Contract No. 33677
Case no SDG H3681

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	% SOLIDS	SAMPLE AMOUNT	BASIS AMOUNT	DAYS S		LAB SAMPLE ID	DEPARTMENT
7075	F08-043-150	B1TH05	SOLID	93.4	129 g		04/09/08	21	R804057-01	7075~001
		Method Blank Lab Control Sample Duplicate (R804057-01)	SOLID SOLID	93.4	129 g		04/09/08	21	R804057-03 R804057-02 R804057-04	7075-003 7075-002 7075-004

QC SUMMARY

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP H3681

SDG 7075
Contact Melissa C. Mannion

PREP BATCH SUMMARY

Client Hanford
Contract No. 33677
Case no SDG H3681

			PREPARATION	ERROR			- PLA	NCHETS A	ANALYZ	ED	- I.JAUQ
TEST	MATRIX	METHOD	BATCH	2o %	CLIENT	MORE	RE	BLANK	LCS	DUP/ORIG MS/ORIG	FIERS
Alpha	Spectroso	сору									
TH	SOLID	Thorium, Isotopic in Solids	6152-064	8.0	1			1	1	1/1	
Beta	Counting										
TC	SOLID	Technetium 99 in Solids	6152-064	13.2	1			1	1	1/1	
Liqui	d Scintill	ation Counting									177110411212
Н	SOLID	Tritium in Solids	6152-064	10.0	1			1	1	1/1	
NI_L	SOLID	Nickel 63 in Solids	6152-064	11.2	1			1	1	1/1	***************************************

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.

Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

PREP BATCH SUMMARY
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SAMPLE DELIVERY GROUP H3681

SDG 7075
Contact Melissa C. Mannion

LAB WORK SUMMARY

Client <u>Hanford</u>
Contract <u>No. 33677</u>
Case no <u>SDG H3681</u>

LAB SAMPLE COLLECTED RECEIVED	CLIENT SAMPLE LOCATION CUSTODY	ID SAF No	MATRIX	PLANCHET	TEST	SUF-	ANALYZED	REVIEWED	ву	METHOD
R804057-01	B1TH05			7075-001	Н		04/26/08	04/29/08	BW	Tritium in Solids
03/19/08	C5941, I-033		SOLID	7075-001	NI_L		05/16/08	05/20/08	BW	Nickel 63 in Solids
04/09/08	F08-043-150	F08-043		7075-001	TC		05/06/08	05/06/08	BW	Technetium 99 in Solids
				7075-001	TH		05/13/08	05/14/08	BW	Thorium, Isotopic in Solids
R804057-02	Lab Control Sa	mple		7075-002	Н		04/26/08	04/29/08	BW	Tritium in Solids
			SOLID	7075-002	NI_L		05/16/08	05/20/08	BW	Nickel 63 in Solids
		F08-043		7075-002	TC		05/03/08	05/06/08	BW	Technetium 99 in Solids
				7075-002	TH		05/13/08	05/14/08	BW	Thorium, Isotopic in Solids
R804057-03	Method Blank			7075-003	Н		04/26/08	04/29/08	BW	Tritium in Solids
			SOLID	7075-003	NI_L		05/16/08	05/20/08	BW	Nickel 63 in Solids
		F08-043		7075-003	TC		05/03/08	05/06/08	BW	Technetium 99 in Solids
				7075-003	TH		05/13/08	05/14/08	BW	Thorium, Isotopic in Solids
R804057-04	Duplicate (R80	4057-01)		7075-004	Н		04/26/08	04/29/08	BW	Tritium in Solids
03/19/08	C5941, I-033		SOLID	7075-004	NI_L		05/16/08	05/20/08	BW	Nickel 63 in Solids
04/09/08		F08-043		7075-004	TC		05/03/08	05/06/08	BW	Technetium 99 in Solids
				7075-004	ТН		05/13/08	05/14/08	BW	Thorium, Isotopic in Solids

		COUNTS OF	TESTS BY	SAMPLE TYPE				
TEST	SAF No	METHOD	REFERENCE	CLIENT MORE	RE BLANK	LCS	DUP SPIKE	TOTAL
Н	F08-043	Tritium in Solids	TRITIUM_COX_LSC	1	1	1	1	4
NI_L	F08-043	Nickel 63 in Solids	NI63_LSC	1	1	1	1	4
TC	F08-043	Technetium 99 in Solids	TC99_TR_SEP_GPC	1	1	1	1	4
тн	F08-043	Thorium, Isotopic in Solids	THISO_IE_PLATE_AEA	1	1	1	1	4
TOTALS				4	4	4	4	16

WORK SUMMARY

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SUMMARY DATA SECTION

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EBERLINE SERVICES/RICHMOND SAMPLE DELIVERY GROUP H3681

7075-003

METHOD BLANK

Method Blank

	7075 Melissa C. Mannion	Client/Case no Contract	Hanford No. 33677	SDG_H3681
 Lab sample id Dept sample id		Client sample id Material/Matrix SAF No		SOLID

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	0	3.6	6.18	400	U	Н
Nickel 63	13981-37-8	0.373	1.7	2.88	30.0	U	NI L
Technetium 99	14133-76-7	-0.007	0.16	0.549	15.0	U	TC
Thorium 228	14274-82-9	0.006	0.098	0.176	1.00	U	TH
Thorium 230	14269-63-7	-0.006	0.11	0.220	1.00	U	TH
Thorium 232	TH-232	-0.018	0.024	0.058	1.00	U	TH

216-A-30 Crib Sampling

QC-BLANK #65344

METHOD BLANKS

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP H3681

7075-002

LAB CONTROL SAMPLE

Lab Control Sample

SDG	7075	Client/Case no	Hanford	SDG H3681
Contact	Melissa C. Mannion	Contract	No. 33677	
Lab sample id	R804057-02	Client sample id	Lab Control Sample	
Dept sample id	7075-002	Material/Matrix		SOLID
		SAF No	F08-043	

ANALYTE	RESULT pCi/g	20 ERR	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST	ADDED pCi/g	2σ ERR pCi/g	REC %	30 LMTS F	PROTOCOL LIMITS
Tritium	843	17	6.86	400		Н	823	33	102	83-117	80-120
Nickel 63	243	6.1	2.94	30.0		NI_L	266	11	91	83-117	80-120
Technetium 99	114	2.6	0.570	15.0		TC	120	4.8	95	80-120	80-120
Thorium 230	17.8	1.0	0.211	1.00		тн	21.9	0.88	81	87-113	80-120

216-A-30 Crib Sampling

QC-LCS #65343		

LAB CONTROL SAMPLES
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SUMMARY DATA SECTION
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Lab id <u>EBRLNE</u>

Protocol <u>Hanford</u>

Version <u>Ver 1.0</u>

Form <u>DVD-LCS</u>

Version <u>3.06</u>

Report date <u>05/22/08</u>

SAMPLE DELIVERY GROUP H3681

7075-004

DUPLICATE

B1TH05

SDG 7075		Client/Case no <u>Hanford</u>	SDG H3681
Contact Melissa C. Mannion		Contract No. 33677	
DUPLICATE	ORIGINAL		
Lab sample id <u>R804057-04</u>	Lab sample id <u>R804057-01</u>	Client sample id B1TH05	
Dept sample id 7075-004	Dept sample id 7075-001	Location/Matrix C5941, I-033	SOLID
	Received 04/09/08	Collected/Weight 03/19/08 08:50	129 g
% solids <u>93.4</u>	% solids <u>93.4</u>	Custody/SAF No F08-043-150 F	08-043

ANALYTE	DUPLICATE pCi/g	20 ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST	ORIGINAL pCi/g	20 ERR	MDA pCi/g	QUALI- FIERS	RPD %	3σ TOT	DER O
Tritium	1.21	2.9	4.88	400	Ü	Н	-0.477	2.9	4.94	U	_		0.8
Nickel 63	0.761	2,1	3.59	30.0	U	NI_L	0.245	2.0	3.46	U	_		0.4
Technetium 99	0.046	0.20	0.638	15.0	Ü	TC	0.194	0.36	0.627	U	-		0.7
Thorium 228	0.439	0.14	0.176	1.00		TH	0.459	0.13	0.119		4	66	0.2
Thorium 230	0.402	0.14	0.211	1.00		TH	0.435	0.15	0.203		8	75	0.3
Thorium 232	0.618	0.13	0.084	1.00		TH	0.474	0.11	0.081		26	50	1.6

216-A-30 Crib Sampling

OC-DUP#1 6	534	5

DUPLICATES

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SUMMARY DATA SECTION

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Lab id EBRLNE
Protocol Hanford
Version Ver 1.0
Form DVD-DUP

Version 3.06

Report date <u>05/22/08</u>

EBERLINE SERVICES / RICHMOND SAMPLE DELIVERY GROUP H3681

7075-001

DATA SHEET

BlTH05

		7075 Melissa C. Mannion	Client/Case no Contract	Hanford No. 33677	SDG H3681
***************************************	Lab sample id Dept sample id Received % solids	7075-001 04/09/08	Client sample id Location/Matrix Collected/Weight Custody/SAF No	C5941, I-033 03/19/08 08:50	SOLID 129 q F08-043

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	-0.477	2.9	4.94	400	U	Н
Nickel 63	13981-37-8	0.245	2.0	3.46	30.0	U	NI L
Technetium 99	14133-76-7	0.194	0.36	0.627	15.0	U	TC
Thorium 228	14274-82-9	0.459	0.13	0.119	1.00		TH
Thorium 230	14269-63-7	0.435	0.15	0.203	1.00		TH
Thorium 232	TH-232	0.474	0.11	0.081	1.00		TH

216-A-30 Crib Sampling

DATA SHEETS
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SAMPLE DELIVERY GROUP H3681

Test TH Matrix SOLID

SDG 7075

Contact Melissa C. Mannion

LAB METHOD SUMMARY

THORIUM, ISOTOPIC IN SOLIDS
ALPHA SPECTROSCOPY

Client Hanford
Contract No. 33677
Contract SDG H3681

RESULTS

Preparation batch	1 6152-064		
R804057-01	7075-001	B1TH05	0.435
R804057-02	7075-002	Lab Control Sample	LOW
R804057-03	7075-003	Method Blank	Ü
R804057-04	7075-004	Duplicate (R804057-01)	ok

METHOD PERFORMANCE

LAB	RAW SUF-	MAX MDA	ALIQ	PREP		YIELD							ANAL-	
SAMPLE ID	TEST FIX CLIENT SAMPLE ID	pCi/g	g	FAC	TION	%	જ	min	keV	KeV	HELD	PREPARED	YZED	DETECTOR
Preparation	batch 6152-064 20 prep error 8	.0 % Re:	ference	Lab N	oteboo	k #6152	pç	g. 64						
R804057-01	B1TH05	0.203	0.250			95		849			55	05/13/08	05/13	SS-033
R804057-02	Lab Control Sample	0.211	0.250			87		849				05/13/08	05/13	SS-034
R804057-03	Method Blank	0.220	0.250			90		846				05/13/08	05/13	SS-035
R804057-04	Duplicate (R804057-01)	0.211	0.250			92		846			55	05/13/08	05/13	SS-036
Nominal val	ues and limits from method	1.00	0.250			20-105	5	150			180			

PROCEDURES	REFERENCE	THISO_IE_PLATE_AEA
	SPP-073	Soil Leaching 10-200 g Aliquot, rev 0
	CP-900	Thorium in Water and Dissolved Solid Samples by
		Extraction Chromatography, rev 1
	CP-008	Heavy Element Electroplating, rev 9

AVERAGES ± 2 SD MDA 0.211 ± 0.014 FOR 4 SAMPLES YIELD 91 ± 7

METHOD SUMMARIES

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP H3681

* Test TC Matrix SOLID
SDG 7075
Contact Melissa C. Mannion

LAB METHOD SUMMARY

TECHNETIUM 99 IN SOLIDS
BETA COUNTING

Client Hanford
Contract No. 33677
Contract SDG H3681

RESULTS

Preparation	batch 6152-064				
R804057-01	7075-00	B1TH05	U		
R804057-02	7075-00	Lab Control Sample	ok		
R804057-03	7075-00	Method Blank	U		
R804057-04	7075-004	Duplicate (R804057-01)	- U		

METHOD PERFORMANCE

LAB SAMPLE ID	RAW SUF- TEST FIX CLIENT SAMPLE ID	MDA pCi/g	ALIQ g	PREP FAC		YIELD	EFF %	COUNT min	 DRIFT KeV	DAYS HELD	PREPARED	ANAL- YZED	DETECTOR
Preparation	batch 6152-064 20 prep error 1	.3,2 % Rei	ference	Lab N	Notebool	c #6152	s ba	j. 64					
R804057-01	B1TH05	0.627	1.00			82		50		48	04/30/08	05/06	GRB-203
R804057-02	Lab Control Sample	0.570	1.00			90		50			04/30/08	05/03	GRB-222
R804057-03	Method Blank	0.549	1.00			94		50			04/30/08	05/03	GRB-223
R804057-04	Duplicate (R804057-01)	0.638	1.00			80		50		45	04/30/08	05/03	GRB-224
Nominal val	ues and limits from method	15.0	1.00			20-105	5	50		180			

	CP-431	Technetium-99 Purification of Soil or Resin by Extraction Chromatography, rev 2
	CP-008	Heavy Element Electroplating, rev 9

AVERAGES ± 2 SD MDA 0.596 ± 0.086 FOR 4 SAMPLES YIELD 86 ± 13

METHOD SUMMARIES

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP H3681

Test H Matrix SOLID
SDG 7075
Contact Melissa C. Mannion

LAB METHOD SUMMARY

TRITIUM IN SOLIDS

Client <u>Hanford</u>
Contract <u>No. 33677</u>
Contract <u>SDG H3681</u>

RESULTS

	SUF-	CLIENT SAMPLE ID	Tritium	
Preparation bate	th 6152-064			
R804057-01	7075-001	B1TH05	U	
R804057-02	7075-002	Lab Control Sample	ok	
R804057-03	7075-003	Method Blank	U	
R804057-04	7075-004	Duplicate (R804057-01)	- Ŭ	

METHOD PERFORMANCE

LAB SAMPLE ID	RAW SUF- TEST FIX CLIENT SAMPLE ID	MDA pCi/q	ALIQ	PREP FAC	DILU- TION	YIELD	EFF %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	PREPARED	ANAL-	DETECTOR
JAME III III			~ · · · · · · · · · · · · · · · · · · ·											
Preparation	batch 6152-064 20 prep error 10).0 % Re	eference	Lab N	iotebool	k #6152	5 be	. 64						
R804057-01	B1TH05	4.94	0.384			100		50			38	04/25/08	04/26	LSC-007
R804057-02	Lab Control Sample	6.86	0.300			100		50				04/25/08	04/26	LSC-007
R804057-03	Method Blank	6.18	0.300			100		50				04/25/08	04/26	LSC-007
R804057-04	Duplicate (R804057-01)	4.88	0.385			100		50			38	04/25/08	04/26	LSC-007
Nominal val	ues and limits from method	400	0.300				****	25			180			

PROCEDURES	REFERENCE	TRITIUM_COX_LSC
	CP-251	Tritium/Carbon-14 Oxidation, rev 8

AVERAGES ± 2 SD	MDA _	5.72	+ 1.94
FOR 4 SAMPLES	YIELD _	100 :	<u>+</u> 0

METHOD SUMMARIES $\begin{array}{ccc} \text{Page} & 3 \\ \\ \text{SUMMARY} & \text{DATA} & \text{SECTION} \\ \\ \text{Page} & 13 \\ \end{array}$

SAMPLE DELIVERY GROUP H3681

Test NI L Matrix SOLID

SDG 7075

Contact Melissa C. Mannion

LAB METHOD SUMMARY

NICKEL 63 IN SOLIDS
LIQUID SCINTILLATION COUNTING

Client Hanford
Contract No. 33677
Contract SDG H3681

RESULTS

LAB	RAW SUF-				
SAMPLE ID	TEST FIX	PLANCHET	CLIENT SAMPLE ID	Nickel	63
Preparation	hatch 615	2 - 064	an alaman an a		Concession and the second
R804057-01	bacch 013	7075-001	B1 T H05	Ū	
R804057-02		7075-002	Lab Control Sample	ok	
R804057-03		7075-003	Method Blank	Ū	
R804057-04		7075-004	Duplicate (R804057-01)	-	U

METHOD PERFORMANCE

LAB	RAW SUF-	MDA	ALIQ	PREP	DILU-	YIELD	EFF	COUNT	FWHM	DRIFT	DAYS		ANAL-	
SAMPLE ID	TEST FIX CLIENT SAMPLE ID	pCi/g	g	FAC	TION	왕	챵	min	keV	KeV	HELD	PREPARED	YZED	DETECTOR
Preparation	batch 6152-064 20 prep error 13	1.2 % Re	ference	Lab 1	Notebool	k #6152	pg s	. 64		######################################	**************************************			The second secon
R804057-01	B1TH05	3.46	0.500			81		50			58	05/14/08	05/16	LSC-004
R804057-02	Lab Control Sample	2.94	0.500			96		50				05/14/08	05/16	LSC-004
R804057-03	Method Blank	2.88	0.500			97		50				05/14/08	05/16	LSC-004
R804057-04	Duplicate (R804057-01)	3.59	0.500			79		50			58	05/14/08	05/16	LSC-004
Nominal val	ues and limits from method	30.0	0.500			30-105	<u> </u>	25		<u> </u>	180	· · · · · · · · · · · · · · · · · · ·		-

ĺ	PROCEDURES	REFERENCE	NI63_LSC
		SPP-071	Soil Dissolution, > 1.0g Aliquot, rev 5
		CP-280	Nickel-63 Purification, rev 3
L			

AVERAGES ± 2 SD	MDA	3.22	±	0.720
FOR 4 SAMPLES	YIELD	88	±	

METHOD SUMMARIES

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SAMPLE DELIVERY GROUP H3681

SDG 7075
Contact Melissa C. Mannion

REPORT GUIDE

Clie	nt	Hanford				
Contra	ct	No.	33677			
Case	no	SDG	H3681			

SAMPLE SUMMARY

The Sample and QC Summary Reports show all samples, including QC samples, reported in one Sample Delivery Group (SDG).

The Sample Summary Report fully identifies client samples and gives the corresponding lab sample identification. The QC Summary Report shows at the sample level how the lab organized the samples into batches and generated QC samples. The Preparation Batch and Method Summary Reports show this at the analysis level.

The following notes apply to these reports:

- * LAB SAMPLE ID is the lab's primary identification for a sample.
- * DEPARTMENT SAMPLE ID is an alternate lab id, for example one assigned by a radiochemistry department in a lab.
- * CLIENT SAMPLE ID is the client's primary identification for a sample. It includes any sample preparation done by the client that is necessary to identify the sample.
- * QC BATCH is a lab assigned code that groups samples to be processed and QCed together. These samples should have similar matrices.

QC BATCH is not necessarily the same as SDG, which reflects samples received and reported together.

* All Lab Control Samples, Method Blanks, Duplicates and Matrix Spikes are shown that QC any of the samples. Due to possible reanalyses, not all results for all these QC samples may be relevant to the SDG. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.

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Lab id <u>EBRLNE</u>

Protocol <u>Hanford</u>

Version <u>Ver 1.0</u>

Form <u>DVD-RG</u>

Version <u>3.06</u>

Report date <u>05/22/08</u>

SAMPLE DELIVERY GROUP H3681

SDG 7075
Contact Melissa C. Mannion

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PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches in one Sample Delivery Group (SDG) with information necessary to check the completeness and consistency of the SDG.

The following notes apply to this report:

- * The preparation batches are shown in the same order as the Method Summary Reports are printed.
- * Only analyses of planchets relevant to the SDG are included.
- * Each preparation batch should have at least one Method Blank and LCS in it to validate client sample results.
- * The QUALIFIERS shown are all qualifiers other than U, J, B, L and H that occur on any analysis in the preparation batch. The Method Summary Report has these qualifiers on a per sample basis.

These qualifiers should be reviewed as follows:

- X Some data has been manually entered or modified. Transcription errors are possible.
- P One or more results are 'preliminary'. The data is not ready for final reporting.
- 2 There were two or more results for one analyte on one planchet imported at one time. The results in DVD may not be the same as on the raw data sheets.

Other lab defined qualifiers may occur. In general, these should be addressed in the SDG narrative.

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WORK SUMMARY

The Work Summary Report shows all samples, including QC samples, and all relevant analyses in one Sample Delivery Group (SDG). This report is often useful as supporting documentation for an invoice.

The following notes apply to this report:

- * TEST is a code for the method used to measure associated analytes. Results and related information for each analyte are on the Data Sheet Report. In special cases, a test code used in the summary data section is not the same as in associated raw data. In this case, both codes are shown on the Work Summary.
- * SUFFIX is the lab's code to distinguish multiple analyses (recounts, reworks, reanalyses) of a fraction of the sample. The suffix indicates which result is being reported. An empty suffix normally identifies the first attempt to analyze the sample.
- * The LAB SAMPLE ID, TEST and SUFFIX uniquely identify all supporting data for a result. The Method Summary Report for each TEST has method performance data, such as yield, for each lab sample id and suffix and procedures used in the method.
- * PLANCHET is an alternate lab identifier for work done for one test. It, combined with the TEST and SUFFIX, may be the best link to raw data.
- * For QC samples, only analyses that directly QC some regular sample are shown. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.
- * The SAS (Special Analytical Services) Number is a client or lab assigned code that reflects special processing for samples, such as rapid turn around. Counts of tests done are lists by SAS number since it is likely to affect prices.

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DATA SHEET

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- * TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- * The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- * ERRORS can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- * A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- * When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

U The RESULT is less than the MDA (Minimum Detectable Activity).

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DATA SHEET

If the MDA is blank, the ERROR is used as the limit.

- J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.
- B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

- L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.
- H Similar to 'L' except the recovery was high.
- P The RESULT is 'preliminary'.
- X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.
- 2 There were two or more results available for this analyte. reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

* An MDA is underlined if it is bigger than its RDL.

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- * An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- * A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- * When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

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LAB CONTROL SAMPLE

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- * An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits for the recovery.
- * The recovery is underlined if it is outside either of these ranges.

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DUPLICATE

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTs divided by their average expressed as a percent.

If both RESULTs are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

* The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTs prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTs. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- * The second limit for the RPD is the larger of:
 - 1. A fixed percentage specified in the protocol.

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- 2. A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.
- * The RPD is underlined if it is greater than either limit.
- * If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

* The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit.

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MATRIX SPIKE

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.
 - If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.
- * An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.
 - An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.
- * REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The errors of the two RESULTs, including those introduced by rounding them prior to printing.
 - If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.
 - 2. The error of ADDED.
 - 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits

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MATRIX SPIKE

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

* The recovery is underlined (out of spec) if it is outside either of these ranges.

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METHOD SUMMARY

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

* Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

* The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

* If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- * Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- * Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

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means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- * Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
- * If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.

MDAs are underlined if greater than the printed RDL.

- * Aliquots are underlined if less than the nominal value specified for the method.
- * Prepareation factors are underlined if greater than the nominal value specified for the method.
- * Dilution factors are underlined if greater than the nominal value specified for the method.
- * Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
- * Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
- * Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

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METHOD SUMMARY

- * Count times are underlined if less than the nominal value specified for the method.
- * Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- * Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- * Days Held are underlined if greater than the holding time specified in the protocol.
- * Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1+3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

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METHOD SUMMARY

results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

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